Contents

No. 1 1-180 issued in September 1988 No. 2 181-364 issued in October 1988

No. 3 365-600 issued in November 1988

Aharonowitz Y → Shiffman D 562 Alonso JC, Stiege CA, Tailor RH, Viret J-F: Functional analysis of the dna (Ts) mutants of Bacillus subtilis: Plasmid pUB110 replication as a model system 482

Altamura S \rightarrow Londei P 48 Altuvia S \rightarrow Suissa M 570 Amils R \rightarrow Londei P 48 Anraku Y \rightarrow Nakao T 379 Asselin A \rightarrow Potvin C 241

Bäckman A → Bailone A 389 Bagdasarian MM → Bailone A 389 Bagdasarian M → Bailone A 389 Bailey JE → Khosla C 158

Bailone A, Bäckman A, Sommer S, Célérier J, Bagdasarian MM, Bagdasarian M, Devoret R: PsiB polypeptide prevents activation of RecA protein in Escherichia coli 389

Bandziulis R, Rosenbaum JL: Novel control elements in the alpha-1 tubulin gene promoter from *Chlamydomonas reinhar*dii 204

Banfalvi Z, Nieuwkoop A, Schell M, Besl L, Stacey G: Regulation of nod gene expression in Bradyrhizobium japonicum 420

420
Baty D → Garcia P 509
Baulcombe DC → Huttly AK 232
Bellemare G → Potvin C 241
Belogurov AA → Efimova EP 313
Belogrov AA → Efimova EP 317
Benner, Lugar D. Boos W: The mall 6:

Benner-Luger D, Boos W: The mgIB sequence of Salmonella typhimurium LT2; promoter analysis by gene fusions and evidence for a divergently oriented gene coding for the mgI repressor 579

Benz R → Ludwig A 553 Besl L → Banfalvi Z 420

Bockrath R, Ruiz-Rubio M: Anti-mutagenic effect of ultraviolet light on spontaneous tyrosine tRNA ochre suppressor mutations in *Escherichia coli* 361

Böhme H, Haselkorn R: Molecular cloning and nucleotide sequence analysis of the gene coding for heterocyst ferredoxin from the cyanobacterium *Anabaena* sp. strain PCC 7120 278

Stant PCC 120 278

Soos W → Benner-Luger D 579

Booth IR → Giffard PM 148

Borstel von RC → Lee GS-F 396

Boulter D → Evans IM 153

Bourgouin C → Delécluse A 42

Bowien B → Husemann M 112

Boynton JE → Liu X-Q 588

Braus G → Paravicini G 165

Brears T, Lonsdale DM: The sugar bee

Brears T, Lonsdale DM: The sugar beet mitochondrial genome: A complex organisation generated by homologous recombination 514 Brehm JK → Oultram JD 177

Brombach M → Friedrich K 595

Bron S → Dijl JM van 55

Bruijn de FJ → Metz BA 181

Bucheton A → Pritchard MA 533

Burcham KWG → Ellis THN 333

Burgess DG, Taylor WC: The chloroplast affects the transcription of a nuclear gene

family 89 Büttcher V → Husemann M 112

Cauvin B → Leclerc M 97
Célérier J → Bailone A 389
Chourey PS, DeRobertis GA, Still PE: Altered tissue specificity of the revertant shrunken allele upon Dissociation (Ds) excision is associated with loss of expression and molecular rearrangement at the corresponding non-allelic isozyme locus in maize 300

Chua N-H → Poulsen C 16 Claverys J-P, Méjean V: Strand targeting signal(s) for in vivo mutation avoidance by post-replication mismatch repair in Escherichia coli 574

Cleary WG → Ellis THN 333 Cohen G → Shiffman D 562 Colbeau A → Leclerc M 97 Crosby RM → Richardson KK 460 Crosby RM → Richardson KK 460

Cseplö A, Etzold T, Schell J, Schreier PH: Point mutations in the 23 S rRNA genes of four lincomycin resistant Nicotiana phumbaginifolia mutants could provide new selectable markers for chloroplast transformation 295

Dankocsik C → Donovan WP 365
Das HK → Raina R 121
Delécluse A, Bourgouin C, Klier A, Rapoport G: Specificity of action on mosquito larvae of *Bacillus thuringiensis israelensis* toxins encoded by two different genes 42

Delver EP → Efimova EP 313 Delver EP → Efimova EP 317

Deng Z, Kieser T, Hopwood DA: "Strong incompatibility" between derivatives of the *Streptomyces* multi-copy plasmid plJ101 286

Dennis ES → Landsmann J 68 DeRobertis GA → Chourey PS 300 Desnoues N → Kaminski PA 496 Devoret R → Bailone A 389

Dick T, Matzura H: Positioning ribosomes on leader mRNA for translational activation of the message of an inducible Staphyloccocus aureus cat gene 108

Dijl JM van, Smith H, Bron S, Venema G: Synthesis and processing of Escherichia coli TEM-β-lactamase and Bacillus licheniformis α-amylase in E. coli: The role of signal peptidase I 55 Doerfler W → Hauser C 373
Donovan WP, Gonzales MJ Jr, Gilbert MP,
Dankocsik C: Isolation and characterization of Ed2158, a new strain of Bacillus
thuringiensis toxic to coleopteran larvae,
and nucleotide sequence of the toxin
gene 365

Doskočil J, Štorchová H, Štokrová J, Forstová J, Meyer J: Correlation of physical maps and some genetic functions in the genomes of the κ-9 phage family of *Bacil*lus licheniformis 343

Drake JW: Bacteriophage T4 DNA polymerase determines the amount and specificity of ultraviolet mutagenesis 547
Drolet M → Echelard Y 503

Dubbert W, Luczak H, Staudenbauer WL: Cloning of two chloramphenicol acetyltransferase genes from Clostridium butyricum and their expression in Escherichia coli and Bacillus subtilis 328

Dura J-M → Pritchard MA 533
Düvell A, Hessberg-Stutzke H, Oeser B,
Rogmann-Backwinkel P, Tudzynski P:
Structural and functional analysis of mitochondrial plasmids in Claviceps purpures 128

Dymetryszyn J → Echelard Y 503

Echelard Y, Dymetryszyn J, Drolet M, Sasarman A: Nucleotide sequence of the hemB gene of Escherichia coli K12 503

Efimova EP, Delver EP, Belogurov AA: Alleviation of type restriction in adenine methylase (dam) mutants of Escherichia coli 313

Efimova EP, Delver EP, Belogurov AA: 2-Aminopurine and 5-bromouracil induce alleviation of type I restriction in Escherichia coli: Mismatches function as inducing signals? 317

Ellis THN, Lee D, Thomas CM, Simpson PR, Cleary WG, Newman M-A, Burcham KWG: 5S rRNA genes in *Pisum*: Sequence, long range and chromosomal organization 333

Elmerich C → Kaminski PA 496 Etzold T → Cseplö A 295

Evans IM, Swinhoe R, Gatehouse LN, Gatehouse JA, Boulter D: Distribution of root mRNA species in other vegetative organs of pea (*Pisum sativum L.*) 153

Finnegan DJ → Pritchard MA 533 Forstová J → Doskočil J 343 Foster PL, Sullivan AD: Interactions between epsilon, the proofreading subunit of DNA polymerase III, and proteins in-

coli 467
Friedrich K, Brombach M, Pon CL: Identification, cloning and sequence of the

volved in the SOS response of Escherichia

Streptococcus faecium infB (translational initiation factor IF2) gene 595

Fu Y-H, Young JL, Marzluf GA: Molecular cloning and characterization of a negative-acting nitrogen regulatory gene of Neurospora crassa 74

Fusswinkel H → Hauser C 373

Ganesan AK, Hunt J, Hanawalt PC: Temperature dependent survival of UV-irradiated Escherichia coli K12 198

Garcia P, Gasc AM, Kyriakidis X, Baty D, Sicard M: DNA sequences required to induce localized conversion in Streptococcus pneumoniae transformation 509

Gasc AM → Garcia P 509 Gatehouse JY → Evans IM 153 Gatehouse LN → Evans IM 153

Gauthier A, Turmel M, Lemieux C: Mapping of chloroplast mutations conferring resistance to antibiotics in Chlamydomonas: Evidence for a novel site of streptomycin resistance in the small subunit rRNA 192

Ghosal D → Raina R 121

Gielow A, Kücherer C, Kölling R, Messer W: Transcription in the region of the replication origin, oriC, of Escherichia coli: Termination of asnC transcripts 474

Giffard PM, Booth IR: The rpo A341 allele of Escherichia coli specifically impairs the transcription of a group of positively-regulated operons 148

Giladi H → Suissa M 570 Gilbert MP → Donovan WP 365 Gillham NW → Liu X-Q 588

Glomp I, Saulnier P, Guespin-Michel J, Schairer HU: Transfer of IncP plasmids into Stigmatella aurantiaca leading to insertional mutants affected in spore development 213

Goebel W → Ludwig A 553 Goldman WE → Worsham PL 348 Golub EI, Panzer HA: The F factor of Escherichia coli carries a locus of stable plasmid inheritance stm, similar to the parB locus of plasmid RI 353

Gomez L, Sanchez-Monge R, Salcedo G: A family of endosperm globulins encoded by genes located in group 1 chromosomes of wheat and related species 541 Gonzales MJ Jr → Donovan WP 365

Gregerson RG → Ortiz DF 135 Guespin-Michel J → Glomp I 213

Hanawalt PC → Ganesan AK 198 Hanks MC, Newman B, Oliver JR, Masters M: Packaging of transducing DNA by bacteriophage P1 523 Harper EC → Maliga P 456

Haselkorn R → Böhme H 278 Hauser C, Fusswinkel H, Li J, Oellig C, Kunze R, Müller-Neumann M, Heinlein M, Starlinger P, Doerfler W: Overproduction of the protein encoded by the maize transposable element Ac in insect

cells by a baculovirus vector 373

Hawkins AR, Lamb HK, Smith M, Keyte JW, Roberts CF: Molecular organisation of the quinic acid utilization (QUT) gene cluster in Aspergillus nidulans 224

Hayashi Y, Kyozuka J, Shimamoto K: Hybrids of rice (Orvza sativa L.) and wild Oryza species obtained by cell fusion 6 Heinlein M → Hauser C 373

Henriquet M → Sibold L 439

Herrmann H, Janke D, Krejsa S, Roy M: In vivo generation of R68.45-pPGH1 hybrid plasmid conferring a Phl+ (meta pathway) phenotype 173

Hessberg-Stutzke H → Düvelt A 128 Hoffmann HJ → Metz BA 181 Hopwood DA → Deng Z 286

Horii T → Tajima Y 451

Hougan L → Whiteway M 85 Hu ST, Lee CH: Characterization of the transposon carrying the STII gene of enterotoxigenic Escherichia coli 490

Hunt J → Ganesan AK 198 Hütter R → Paravicini G 165

Husemann M, Klintworth R, Büttcher V, Salnikow J, Weissenborn C, Bowien B: Chromosomally and plasmid-encoded gene clusters for CO2 fixation (cfx genes) in Alcaligenes eutrophus 112

Huttly AK, Martienssen RA, Baulcombe DC: Sequence heterogeneity and differential expression of the α-Amy2 gene family in wheat 232

lino T → Kutsukake K 11 Ikeda H → Saing KM 1 Itayama H → Tanda S 405 Itoh K → Mizuno T 249 Itoh T → Tajima Y 451

Janke D → Herrmann H 173 Jarchau T → Ludwig A 553 Jensen EØ → Metz BA 181 Jensen SE → Shiffman D 562 Jessberg-Stutzke H → Düvell A 128 Jones JDG → Maliga P 456

Kaminski PA, Norel F, Desnoues N, Kush A, Salzano G, Elmerich C: Characterization of the fixABC region of Azorhizobium caulinodans ORS571 and identification of a new nitrogen fixation gene 496 Karmazyn-Campelli C → Sanchez-Rivas C

Kato M → Muramatsu S 433

Kelly R, Miller SM, Kurtz MB: One-step gene disruption by cotransformation to isolate double auxotrophs in Candida albicans 24

Keyte JW → Hawkins AR 224

Khosla C, Bailey JE: The Vitreoscilla hemoglobin gene: Molecular cloning, nucleotide sequence and genetic expression in Escherichia coli 158

Kieser T → Deng Z 286 Klier A → Delécluse A 42

Klintworth R → Husemann M 112

Koby S → Suissa M 570

Kohara Y → Muramatsu S 433

Kölling R → Gielow A 474 Kreisa S → Herrmann H 173

Kücherer C → Gielow A 474 Küntzel H → Munder T 271

Kunze R, Starlinger P, Schwartz D: DNA methylation of the maize transposable element Ac interferes with its transcription 325

Kunze R → Hauser C 373

Kurtz MB → Kelly R 24

Kush A → Kaminski PA 496

Kutsukake K, Ohya Y, Yamaguchi S, Iino T: Operon structure of flagellar genes in Salmonella typhimurium 11

Kyozuka J → Hayashi Y 6 Kyriakidis X → Garcia P 509

Lamb HK → Hawkins AR 224

Landsmann J, Llewellyn D, Dennis ES, Peacock WJ: Organ regulated expression of the Parasponia andersonii haemoglobin gene in transgenic tobacco plants 68

Langley CH → Tanda S 405

Leclerc D → Potvin C 241 Leclerc M, Colbeau A, Cauvin B, Vignais PM: Cloning and sequencing of the genes encoding the large and the small subunits of the H2 uptake hydrogenase (hup) of Rhodobacter capsulatus 97

Lee CH → Hu ST 490

Lee D → Ellis THN 333

Lee GS-F, Savage EA, Ritzel RG, Borstel von RC: The base-alteration spectrum of spontaneous and ultraviolet radiation-induced forward mutations in the URA3 locus of Saccharomyces cerevisiae 396

Lemieux B → Turmel M 412

Lemieux C → Gauthier A 192 Lemieux C → Turmel M 412

Levi-Meyrueis C → Sanchez-Rivas C 321

Li J → Hauser C 373

Ling-Ling C → Tanda S 405 Lisowsky T, Michaelis G: A nuclear gene essential for mitochondrial replication suppresses a defect of mitochondrial transcription in Saccharomyces cerevisiae 218

Liu X-Q, Gillham NW, Boynton JE: Chloroplast ribosomal protein L-18 in Chlamydomonas reinhardtii is processed during ribosome assembly 588

Llewellyn D → Landsmann J 68 Londei P, Altamura S, Sanz JL, Amils R: Aminoglycoside-induced mistranslation in thermophilic archaebacteria 48

Lonsdale DM → Brears T 514 Loppes R → Matagne RF 257

Luczak H → Dubbert W 328 Ludwig A, Jarchau T, Benz R, Goebel W: The repeat domain of Escherichia coli haemolysin (HlyA) is responsible for its Ca2+-dependent binding to erythrocytes 553

Makris JC → Nordmann PL 62 Maliga P, Svab Z, Harper EC, Jones JDG:

Improved expression of streptomycin resistance in plants due to a deletion in the streptomycin phosphotransferase coding sequence 456

Martienssen RA → Huttly AK 232 Marzluf GA → Fu Y-H 74

Masters M → Hanks MC 523

Matagne RF, Rongvaux D, Loppes R: Transmission of mitochondrial DNA in crosses involving diploid gametes homozygous or heterozygous for the matingtype locus in *Chlamydomonas* 257
Matsubayashi H → Tanda S 405
Matzura H → Dick T 108
Méjean V → Claverys JP 574
Mellemare G → Potvin C 241
Messer W → Gielow A 474
Metz BA, Welters P, Hoffmann HJ, Jensen EØ, Schell J, Bruijn de FJ: Primary structure and promoter analysis of leghemoglobin genes of the stem-nodulated tropical legume *Sesbania rostrata*: Conserved coding sequences, *cis*-elements and *trans*-acting factors 181

Mevarech M → Shiffman D 562 Meyer J → Doskočil J 343 Michaelis G → Lisowsky T 218 Michaelis U, Schlapp T, Rödel G: Yeast nuclear gene CBS2, required for translational activation of cytochrome b, encodes a basic protein of 45 kDa 263 Miller SM → Kelly R 24

Mink $M \rightarrow$ Munder T 271 Minton NP \rightarrow Oultram JD 177 Miura A \rightarrow Saing KM 1 Mizuno T, Itoh K: Random cloning of bent

Mizuno T, Itoh K: Random cloning of bent DNA segments from Saccharomyces cerevisiae and primary characterization of their structures 249

Mizuno T → Muramatsu S 433 Moore PD → Simon JR 37 Müller-Neumann M → Hauser C 373 Munder T, Mink M, Küntzel H: Domains of the Saccharomyces cerevisiae CDC25 gene controlling mitosis and meiosis 271

Muramatsu S, Kato M, Kohara Y, Mizuno T: Insertion sequence IS5 contains a sharply curved DNA structure at its terminus 433

Nakao T, Yamato I, Anraku Y: Mapping of the multiple regulatory sites for putP and putA expression in the putC region of Escherichia coli 379

Newman B → Hanks MC 523

Newman M-A → Ellis THN 333

Nieuwkoop A → Banfalvi Z 420

Nordmann PL, Makris JC, Reznikoff WS: Inosine induced mutations 62

Norel F → Kaminski PA 496

Oellig C → Hauser C 373
Oeser B → Düvell A 128
Ohya Y → Kutsukake K 11
Oliver JR → Hanks MC 523
Oppenheim AB → Suissa M 570
Orii H → Saing KM 1
Ortiz DF, Rowland LJ, Gregerson RG,
Strommer JN: Insertion of Mu into the
Shrunken 1 gene of maize affects transcriptional and post-transcriptional regulation of Sh1 RNA 135
Oultram JD, Peck H, Brehm JK, Thompson

Dultram JD, Peck H, Brenm JK, I nompson DE, Swinfield TJ, Minton NP: Introduction of genes for leucine biosynthesis from Clostridium pasteurianum into C. acetobutylicum by cointegrate conjugal transfer 177

Palva ET → Pirhonen M 170 Panzer HA → Golub El 353 Paravicini G, Braus G, Hütter R: Structure of the ARO3 gene of Saccharomyces cerevisiae 165 Peacock WJ → Landsmann J 68 Peck H → Oultram JD 177

Pélisson A → Pritchard MA 533

Pirhonen M, Palva ET: Occurence of bacteriophage T4 receptor in *Erwinia caroto*vora 170

Piškur J: Transmission of yeast mitochondrial loci to progeny is reduced when nearby intergenic regions containing ori sequences are deleted 425

Pon CL → Friedrich K 595
Potvin C, Leclere D, Tremblay G, Asselin
A, Bellemare G: Cloning, sequencing and
expression of a Bacillus bacteriolytic enzyme in Escherichia coli 241

Poulsen C, Chua N-H: Dissection of 5' upstream sequences for selective expression of the Nicotiana plumbaginifolia rbcS-8B gene 16

Pritchard MA, Dura J-M, Pélisson A, Bucheton A, Finnegan DJ: A cloned I-factor is fully functional in *Drosophila melano*gaster 533

Raina R, Reddy MA, Ghosal D, Das HK:
Characterization of the gene for the Feprotein of the vanadium dependent alternative nitrogenase of Azotobacter vinelandii and construction of a Tn5 mutant 121
Rapoport G→ Delécluse A 42

Reddy MA → Raina R 121
Reznikoff WS → Nordmann PL 62
Richardson KK, Crosby RM, Skopek TR:
Mutation spectra of N-ethyl-N'-nitro-Nnitrosoguanidine and 1-(2-hydroxyethyl)1-nitrosourea in Escherichia coli 460
Ritzel RG → Lee GS-F 396

Roberts CF → Hawkins AR 224 Rödel G → Michaelis U 263 Rogmann-Backwinkel P → Düvell A 128 Rongvaux D → Matagne RF 257 Rosenbaum JL → Bandziulis R 204 Rowland LJ → Ortiz DF 135 Roy M → Herrmann H 173

Ruiz-Rubio M → Bockrath R 361

Saigo K → Tanda S 405
Saing KM, Orii H, Tanaka Y, Yanagisawa K, Miura A, Ikeda H: Formation of deletion in *Escherichia coli* between direct repeats located in the long inverted repeats of a cellu'ar slime mold plasmid: Participation of DNA gyrase 1
Salcedo G → Gomez L 541
Salnikow J → Husemann M 112
Salzano G → Kaminski PA 496

Salnikow J → Husemann M 112
Salzano G → Kaminski PA 496
Sanchez-Monge R → Gomez L 541
Sanchez-Rivas C, Karmazyn-Campelli C,
Levi-Meyrueis C: Further studies on recombination in dipliod clones from *Bacil-*lus subtilis protoplast fusion 321
Sanz JL → Londei P 48

Saniar A → Echelard Y 503
Sato F, Shigematsu Y, Yamada Y: Selection of an atrazine-resistant tobacco cell line having a mutant psbA gene 358
Saulnier P → Glomp I 213
Savage EA → Lee GS-F 396

Schell J → Cseplö A 295
Schell J → Metz BA 181
Schell M → Banfalvi Z 420
Schlapp T → Michaelis U 263
Schreier PH → Cseplö A 295
Schwartz D → Kunze R 325
Sherratt DJ → Stirling CJ 80
Shiffman D, Mevarech M, Jensen SE, Cohen G, Aharonowitz Y: Cloning and comparative sequence analysis of the ge coding for isopenicillin N synthase in

Schairer HU → Glomp I 213

hen G, Aharonowitz Y: Cloning and comparative sequence analysis of the gene coding for isopenicillin N synthase in Streptomyces 562
Shigematsu Y → Sato F 358
Shimamoto K → Hayashi Y 6

Shimamoto K → Hayashi Y 6
Shrimpton AE → Tanda S 405
Sibold L, Henriquet M: Cloning of the trp
genes from the archaebacterium Methanococcus voltae: Nucleotide sequence of
the trpBA genes 439

Sicard M → Garcia P 509 Simon JR, Moore PD: Induction of homologous recombination in Saccharomyces cerevisiae 37

Simpson PR → Ellis THN 333
Skopek TR → Richardson KK 460
Smith H → Dijl JM van 55
Smith M → Hawkins AR 224
Sommer S → Bailone A 389
Stacey G → Banfalvi Z 420
Starlinger P → Hauser C 373
Starlinger P → Kunze R 325
Staudenbauer WL → Dubbert W 328

Staudenbauer WL → Dubbert W 328 Stauffer GV, Stauffer LT: Salmonella typhimurium LT2metF operator mutations 32 Stauffer LT → Stauffer GV 32

Stauffer LT → Stauffer GV 32 Stewart G → Stirling CJ 80 Stiege CA → Alonso JC 482 Still PE → Chourey PS 300

Stirling CJ, Stewart G, Sherratt DJ: Multicopy plasmid stability in *Escherichia coli* requires host-encoded functions that lead to plasmid site-specific recombination 80

Štokrová J → Doskočil J 343 Štorchová H → Doskočil J 343 Strommer JN → Ortiz DF 135

Suissa M, Altuvia S, Koby S, Giladi H, Oppenheim AB: Translational signals of a major head protein gene of bacteriophage lambda 570

Sullivan AD → Foster PL 467 Svab Z → Maliga P 456 Swinfield TJ → Oultram JD 177 Swinhoe R → Evans IM 153

Tailor RH → Alonso JC 482
Tajima Y, Horii T, Itoh T: Replication of CoIE2 and CoIE3 plasmids: Two CoIE2 incompatibility functions 451
Tanaka K → Toh-e A 162
Tanaka Y → Saing KM 1
Tanda S, Shrimpton AE, Ling-Ling C
Itayama H, Matsubayashi H, Saigo K,
Tobari YN, Langley CH: Retrovirus-like features and site specific insertions of a transposable element, tom, in Drosophila ananassae 405
Taylor WC → Burgess DG 89
Thomas CM → Ellis THN 333

Thomas CM → Ellis THN 333 Thomas DY → Whiteway M 85 Thompson DE → Oultram JD 177 Thomson J → Webster J 142
Tobari YN → Tanda S 405
Tohe A, Tanaka K, Uesono Y, Wickner
RB: PHO85, a negative regulator of the
PHO system, is a homolog of the protein
kinase gene, CDC28, of Saccharomyces

cerevisiae 162

Tremblay G → Potvin C 241

Tudzynski P → Düvell A 128

Turmel M, Lemieux B, Lemieux C: The chloroplast genome of the green alga
Chlamydomonas moewusii: Localization of protein-coding genes and transcriptionally active regions 412

Turmel M → Gauthier A 192
Uesono Y → Toh-e A 162

Venema G → Dijl JM van 55 Vignais PM → Leclerc M 97 Viret J-F → Alonso JC 482

Waleh NS: Functional expression of Aquaspirillum magnetotacticum genes in Escherichia coli K12 592

Webster J, Thomson J: Genetic analysis of an Agrobacterium tumefaciens strain producing an agrocin active against biotype 3 pathogens 142

Weissenborn C → Husemann M 112

Welters P → Metz BA 181

Whiteway M, Hougan L, Thomas DY: Expression of MFαI in MATa cells supersensitive to α-factor leads to self-arrest 85

Wickner RB → Toh-e A 162
Worsham PL, Goldman WE: Selection and characterization of *ura5* mutants of *Histoplasma capsulatum* 348

Xiao W: Stability of cloned Brassica napus chloroplast DNA fragments in the cyanobacterium Anacystis nidulans R2 307

Yamada Y → Sato F 358 Yamaguchi S → Kutsukake K 11 Yamato I → Nakao T 379 Yanagisawa K → Saing KM 1 Young JL → Fu Y-H 74

Erratum 180

Indexed in Current Contents

